

EAST Search History

Ref #	Hits	Search Query	DBs	Default Operator	Plurals	Time Stamp
S1	1	10/719311	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2006/06/15 11:26
S2	18	Chiorini John	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:18
S3	21	AAV4.clm.	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:24
S4	12	Safer Brian	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:22
S5	26	Kotin Robert	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:21
S6	32	S2 S4 S5	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2006/06/15 11:21
S7	8	S6 and S3	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2006/06/15 11:21
S8	18	Chiorini John	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:22
S9	96	AAV4	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:24
S10	9	S9 and zhang	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:25
S11	2	("6194191").PN.	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT	OR	OFF	2006/06/15 11:26
S12	14214	adeno-associated virus	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:27
S15	4068	S12 and (Rep OR capsid)	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:28

EAST Search History

S17	428	S12 and (Rep capsid ITR)	US-PGPUB; USPAT; EPO; JPO; DERWENT	SAME	ON	2006/06/15 11:36
S18	114	S17 and (AAV2 AAV3 AAV4 AAV5)	US-PGPUB; USPAT; EPO; JPO; DERWENT	OR	ON	2006/06/15 11:56
S19	298	Johnson Philip	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:56
S20	18	S19 and S12	US-PGPUB; USPAT; EPO; JPO; DERWENT	NEAR	ON	2006/06/15 11:56

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(FILE 'HOME' ENTERED AT 16:32:20 ON 22 JUN 2006)

FILE 'MEDLINE, SCISEARCH, CAPLUS, BIOSIS' ENTERED AT 16:34:35 ON 22 JUN 2006

L1 89 S AAV4 OR (ADENO? VIRUS TYPE 4)
L2 34 DUP REM L1 (55 DUPLICATES REMOVED)
L3 5 S L2 AND PY<=1997
L4 5 SORT L3 PY
L5 17 S L2 AND (REP OR CAP? OR ITR?)
L6 17 FOCUS L5 1-
E CHIORINI JOHN?/AU
L7 125 S E1
E SAFER BRAIN?/AU
L8 372 S E1
E KOTIN ROBERT?/AU
L9 114 S E1
L10 574 S L7 OR L8 OR L9
L11 29 S L10 AND L1
L12 13 DUP REM L11 (16 DUPLICATES REMOVED)

=> d ti so au ab pi l6 1 4 6 9

L6 ANSWER 1 OF 17 CAPLUS COPYRIGHT 2006 ACS on STN
TI Adeno-associated virus 4 genome sequence and uses as a genetic vector
SO PCT Int. Appl., 80 pp.
CODEN: PIXXD2
IN Chiorini, John A.; Kotin, Robert M.; Safer, Brian
AB The present invention provides an adeno-associated virus 4 (AAV4) virus and vectors and particles derived therefrom. To understand the nature of AAV4 virus and to determine its usefulness as a vector for gene transfer, it was cloned and sequenced. AAV4 is a distinct virus based on sequence anal. phys. properties of the virion, hemagglutination activity, and tissue tropism. The sequence data indicates that AAV4 is a distinct virus from that of AAV2. In contrast to original reports, AAV4 contains 2 open reading frames which code for either Rep proteins or capsid proteins. AAV4 contains addnl. sequence upstream of the p5 promoter which may affect promoter activity, packaging, or particle stability. Furthermore, AAV4 contains an expanded Rep binding site in its ITR which could alter its activity as an origin of replication or promoter. In contrast to previous reports AAV4 is able to transduce human as well as monkey cells. The inverted terminal repeats may be used to construct vectors containing a promoter and heterologous gene.

	PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
PI	WO 9811244	A2	19980319	WO 1997-US16266	19970911
	W: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
	RW: GH, KE, LS, MW, SD, SZ, UG, ZW, AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG				
	CA 2265460	AA	19980319	CA 1997-2265460	19970911
	AU 9746456	A1	19980402	AU 1997-46456	19970911
	EP 932694	A2	19990804	EP 1997-945204	19970911
	R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO				
	US 2003215422	A1	20031120	US 1999-254747	19991126
	AU 771545	B2	20040325	AU 2001-97210	20011212

AU 2001097210	A5	20020207		
US 2004086490	A1	20040506	US 2003-719311	20031120

L6 ANSWER 4 OF 17 MEDLINE on STN

TI Cloning of adeno-associated virus type 4 (AAV4) and generation of recombinant AAV4 particles.

SO Journal of virology, (1997 Sep) Vol. 71, No. 9, pp. 6823-33.
Journal code: 0113724. ISSN: 0022-538X.

AU Chiorini J A; Yang L; Liu Y; Safer B; Kotin R M

AB We have cloned and characterized the full-length genome of adeno-associated virus type 4 (AAV4). The genome of AAV4 is 4,767 nucleotides in length and contains an expanded p5 promoter region compared to AAV2 and AAV3. Within the inverted terminal repeat (ITR), several base changes were identified with respect to AAV2. However, these changes did not affect the ability of this region to fold into a hairpin structure. Within the ITR, the terminal resolution site and Rep binding sites were conserved; however, the Rep binding site was expanded from three GAGC repeats to four. The Rep gene product of AAV4 shows greater than 90% homology to the Rep products of serotypes 2 and 3, with none of the changes occurring in regions which had previously been shown to affect the known functions of Rep68 or Rep78. Most of the differences in the capsid proteins lie in regions which are thought to be on the exterior surface of the viral capsid. It is these unique regions which are most likely to be responsible for the lack of cross-reacting antibodies and the altered tissue tropism compared to AAV2. The results of our studies, performed with a recombinant version of AAV4 carrying a lacZ reporter gene, suggest that AAV4 can transduce human, monkey, and rat cells. Furthermore, comparison of transduction efficiencies in a number of cell lines, competition cotransduction experiments, and the effect of trypsin on transduction efficiency all suggest that the cellular receptor for AAV4 is distinct from that of AAV2.

L6 ANSWER 6 OF 17 MEDLINE on STN

TI Structure of adeno-associated virus type 4.

SO Journal of virology, (2005 Apr) Vol. 79, No. 8, pp. 5047-58.
Journal code: 0113724. ISSN: 0022-538X.

AU Padron Eric; Bowman Valorie; Kaludov Nikola; Govindasamy Lakshmanan; Levy Hazel; Nick Phillip; McKenna Robert; Muzyczka Nicholas; Chiorini John A; Baker Timothy S; Agbandje-McKenna Mavis

AB Adeno-associated virus (AAV) is a member of the Parvoviridae, belonging to the Dependovirus genus. Currently, several distinct isolates of AAV are in development for use in human gene therapy applications due to their ability to transduce different target cells. The need to manipulate AAV capsids for specific tissue delivery has generated interest in understanding their capsid structures. The structure of AAV type 4 (AAV4), one of the most antigenically distinct serotypes, was determined to 13-A resolution by cryo-electron microscopy and image reconstruction. A pseudoatomic model was built for the AAV4 capsid by use of a structure-based sequence alignment of its major capsid protein, VP3, with that of AAV2, to which AAV4 is 58% identical and constrained by its reconstructed density envelope. The model showed variations in the surface loops that may account for the differences in receptor binding and antigenicity between AAV2 and AAV4. The AAV4 capsid surface topology also shows an unpredicted structural similarity to that of Aleutian mink disease virus and human parvovirus B19, autonomous members of the genus, despite limited sequence homology.

L6 ANSWER 9 OF 17 CAPLUS COPYRIGHT 2006 ACS on STN

TI Recombinant adeno-associated virus type 2, 4, and 5 vectors: transduction of variant cell types and regions in the mammalian central nervous system

SO Proceedings of the National Academy of Sciences of the United States of

America (2000), 97(7), 3428-3432

CODEN: PNASA6; ISSN: 0027-8424

AU Davidson, Beverly L.; Stein, Colleen S.; Heth, Jason A.; Martins, Ines;
Kotin, Robert M.; Derksen, Todd A.; Zabner, Joseph; Ghodsi, Abdi;
Chiorini, John A.

AB Recombinant adeno-associated virus vectors based on serotype 2 (rAAV2) can direct transgene expression in the central nervous system (CNS), but it is not known how other rAAV serotypes perform as CNS gene transfer vectors. Serotypes 4 and 5 are distinct from rAAV2 and from each other in their capsid regions, suggesting that they may direct binding and entry into different cell types. In this study, we examined the tropisms and transduction efficiencies of β -galactosidase-encoding vectors made from rAAV4 and rAAV5 compared with similarly designed rAAV2-based vectors. Injection of rAAV5 β -galactosidase (β gal) or rAAV4 β gal into the lateral ventricle resulted in stable transduction of ependymal cells, with approx. 10-fold more pos. cells than in mice injected with rAAV2 β gal. Major differences between the three vectors were revealed upon striatal injections. Intrastratial injection of rAAV4 β gal resulted again in striking ependyma-specific expression of transgene, with a notable absence of transduced cells in the parenchyma. RAAV2 β gal and rAAV5 β gal intrastratial injections led to β -gal-pos. parenchymal cells, but, unlike rAAV2 β gal, rAAV5 β gal transduced both neurons and astrocytes. The number of transgene-pos. cells in rAAV5 β gal-injected brains was 130 and 5,000 times higher than in rAAV2 β gal-injected brains at 3 and 15 wk, resp. Moreover, transgene-pos. cells were widely dispersed throughout the injected hemisphere in rAAV5 β gal-transduced animals. Together, our data provide in vivo support for earlier in vitro work, suggesting that rAAV4 and rAAV5 gain cell entry by means of receptors distinct from rAAV2. These differences could be exploited to improve gene therapy for CNS disorders.

Kaushal, Sumesh

To: STIC-Biotech/ChemLib
Subject: 10719311: SEQ search

10719311: SEQ search

Please search

	<u>SIZE</u>
• SEQ ID NO: 6	125nt
• SEQ ID NO: 20	129nt
• DNA encoding SEQ ID NO:2	623aa
• DNA encoding SEQ ID NO:8	313aa
• DNA encoding SEQ ID NO:9	399aa
• DNA encoding SEQ ID NO:10	537aa
• DNA encoding SEQ ID NO:11	623aa
• DNA encoding SEQ ID NO:4	734aa
• DNA encoding SEQ ID NO:16	598aa
• DNA encoding SEQ ID NO:18	544aa

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